

=> d bib ab ind 1

LG ANSWER 1 OF 2 CA COPYRIGHT 2003 ACS  
AN 94:188433 CA  
TI Regulation of the basal and cyclic AMP-stimulated rates of glycogen  
synthesis in Escherichia coli by an intermediate of purine biosynthesis  
AU **Leckie, Mary P.**; Porter, Sharon E.; Tieber, Virginia L.;  
Dietzler, David N.  
CS Sch. Med., Washington Univ., St. Louis, MO, 63110, USA  
SO Biochemical and Biophysical Research Communications (1981),  
99(4), 1433-42  
CODEN: BBRCA9; ISSN: 0006-291X  
DT Journal  
LA English  
AB In E. coli, an abrupt increase in the rate of glycogen synthesis occurs at  
the onset of total N starvation. This increase occurs because of the loss  
of a N-contg. intermediate of purine biosynthesis (apparently,  
5-aminoimidazole-4-carboxamide ribonucleotide) that inhibits glycogen  
synthesis. This inhibitory intermediate antagonizes the stimulation of  
glycogen synthesis by cAMP. The regulation of glycogen synthesis by this  
inhibitor provides the 1st link in understanding the reciprocal relation  
between growth rate and glycogen accumulation in E. coli.  
CC 10-2 (Microbial Biochemistry)  
ST glycogen formation regulation Escherichia  
IT Escherichia coli  
(glycogen formation by, regulation of, by purine synthesis  
intermediate)  
IT 9005-79-2, biological studies  
RL: FORM (Formation, nonpreparative)  
(formation of, by Escherichia coli, regulation of, by purine synthesis  
intermediate)  
IT 3031-94-5  
RL: BIOL (Biological study)  
(glycogen formation by Escherichia coli regulation by)  
IT 60-92-4  
RL: BIOL (Biological study)  
(glycogen formation by Escherichia coli regulation by, inhibition of,  
by purine synthesis intermediate)  
IT 9027-71-8  
RL: BIOL (Biological study)  
(of Escherichia coli, purine synthesis intermediates effect on)

L10 ANSWER 8 OF 9 CA COPYRIGHT 2003 ACS

AN 98:155727 CA

TI The in vitro expression of the gene for Escherichia coli ADP-glucose pyrophosphorylase is stimulated by cyclic AMP and cyclic AMP receptor protein

AU Urbanowski, Joseph; Leung, Patrick; Weissbach, Herbert; **Preiss, Jack**

CS Roche Inst. Mol. Biol., Nutley, NJ, 07110, USA

SO Journal of Biological Chemistry (1983), 258(5), 2782-4

CODEN: JBCHA3; ISSN: 0021-9258

DT Journal

LA English

AB The in vitro expression of glgC, the structural gene for E. coli ADP-glucose pyrophosphorylase [9027-71-8], was studied by measuring N-terminal dipeptide and tripeptide synthesis with plasmid pOP12, which carries the glgC gene. When cAMP [60-92-4] and cAMP receptor protein were added together, synthesis of fMet-Val and fMet-Val-Ser was stimulated .apprx.2.5-4-fold. The above tripeptide sequence is the known N-terminal sequence of the ADP-glucose pyrophosphorylase. Hence, cAMP and cAMP receptor protein might be involved in the genetic regulation of the expression of the glgC gene.

L10 ANSWER 6 OF 9 CA COPYRIGHT 2003 ACS  
AN 99:154999 CA  
TI Pyrophosphate may be involved in regulation of bacterial glycogen synthesis  
AU **Preiss, Jack**; Greenberg, Elaine  
CS Dep. Biochem. Biophys., Univ. California, Davis, CA, 95616, USA  
SO Biochemical and Biophysical Research Communications (1983), 115(3), 820-6  
CODEN: BBRCA9; ISSN: 0006-291X  
DT Journal  
LA English  
AB Inorg. pyrophosphate (PPi) is a potent inhibitor of the enzyme that catalyzes synthesis of the glucosyl donor for Escherichia coli glycogen synthesis, ADP-glucose pyrophosphorylase. The  $K_i$  is detd. to be 40  $\mu$ M. ATP, fructose 1,6-diphosphate, and 5'-AMP do not greatly affect the inhibition. PPi exhibits mixed-type inhibition with the other substrate, glucose 1-phosphate. The potential regulation of glycogen synthesis by PPi is discussed.

L13 ANSWER 8 OF 11 CA COPYRIGHT 2003 ACS  
AN 101:106377 CA  
TI Identification of GTP as a physiologically relevant **inhibitor** of  
Escherichia coli **ADP-glucose synthetase**  
AU Dietzler, David N.; Porter, Sharon E.; Roth, William G.; Leckie, Mary P.  
CS Sch. Med., Washington Univ., St. Louis, MO, 63110, USA  
SO Biochemical and Biophysical Research Communications (1984), 122(1), 289-96  
CODEN: BBRCA9; ISSN: 0006-291X  
DT Journal  
LA English  
AB Physiol. concns. of GTP can significantly inhibit wild-type E. coli  
**ADP-glucose synthetase** (the rate-limiting  
enzyme of bacterial glycogen synthesis); mutant-strain enzymes, known to  
show less inhibition by physiol. AMP levels, also show less inhibition by  
physiol. levels of GTP. This decreased inhibition by both AMP and GTP can  
almost totally account for the higher cellular rates of glycogen synthesis  
obsd. in the mutant strains. In addn., in metabolic conditions where  
cellular glycogen synthesis increases, cellular GTP levels are known to  
decrease. Thus, GTP inhibition is physiol. relevant.

=> d

L1 ANSWER 1 OF 1 REGISTRY COPYRIGHT 2003 ACS  
RN 9027-71-8 REGISTRY  
CN Adenylyltransferase, glucose 1-phosphate (9CI) (CA INDEX NAME)  
OTHER NAMES:  
CN Adenosine diphosphate glucose pyrophosphorylase  
CN Adenosine diphosphoglucose pyrophosphorylase  
CN ADP-glucose pyrophosphorylase  
CN ADP-glucose synthase  
CN ADP-glucose synthetase  
CN ADP:.alpha.-D-glucose-1-phosphate adenylyltransferase  
CN ADPG pyrophosphorylase  
CN **E.C. 2.7.7.27**  
CN Glucose 1-phosphate adenylyltransferase  
MF Unspecified  
CI MAN  
LC STN Files: AGRICOLA, ANABSTR, BIOBUSINESS, BIOSIS, BIOTECHNO, CA,  
CAPLUS, CEN, CIN, EMBASE, TOXCENTER, USPAT2, USPATFULL

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

733 REFERENCES IN FILE CA (1957 TO DATE)

1 REFERENCES TO NON-SPECIFIC DERIVATIVES IN FILE CA

734 REFERENCES IN FILE CAPLUS (1957 TO DATE)

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L8 ANSWER 52 OF 59 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.

AN 1980:150/59 BIOSIS

DN BA69:25755

TI EVIDENCE FOR NEW FACTORS IN THE COORDINATED REGULATION OF ENERGY METABOLISM IN ESCHERICHIA-COLI EFFECTS OF HYPOXIA CHLORAMPHENICOL SUCCINATE AND 2 4 DI NITRO PHENOL ON GLUCOSE UTILIZATION GLYCOGEN SYNTHESIS ADENYLATE ENERGY CHARGE AND HEXOSE PHOSPHATES DURING THE 1ST 2 PERIODS OF NITROGEN STARVATION.

AU DIETZLER D N; LECKIE M P; LEWIS J W; PORTER S E; TAXMAN T L; LAIS C J

CS DEP. PEDIATR., WASH. UNIV. SCH. MED., ST. LOUIS, MO. 63110, USA.

SO J BIOL CHEM, (1979) 254 (17), 8295-8307.

CODEN: JBCHA3. ISSN: 0021-9258.

FS BA; OLD

LA English

AB The effects of decreased aeration, chloramphenicol succinate, and 2,4-dinitrophenol on the cellular rates of glycogen synthesis and glucose utilization and on the cellular concentrations of adenine nucleotides, glucose 6-phosphate, fructose 1,6-diphosphate, and phosphoenolpyruvate during the 1st 2 periods of N starvation of *E. coli* W4597(K) were studied. A quantitative relationship between the changes in the rates and the accompanying changes in the hexose phosphates was demonstrated. The relationship for glycogen synthesis was different in different sets of metabolic conditions. This difference reflects a change in the steady state level of a previously unknown effector of **ADP-glucose synthetase** (glucose 1-phosphate adenylyltransferase, EC 2.7.7.27), the rate-limiting enzyme of bacterial glycogen synthesis. The properties of the hypothetical in vivo effector are consistent with the **inhibitory** effects of pppGpp (guanosine 3'-diphosphate 5'-diphosphate) and ppGpp (guanosine 3'-diphosphate 5'-triphosphate) on this enzyme in vitro. Tetracycline, an **inhibitor** of the synthesis of these nucleotides, apparently prevents the change in the quantitative relationship. The relationship between glucose utilization and the hexose phosphates is altered at the transition to period II of N starvation. This change reflects the alteration of the cellular steady state level of an unknown effector of the glucose phosphotransferase system. Unlike the ATP-hexose phosphate system of shared regulatory effects, the specific effects of the unknown effectors allow the rates of glucose utilization and glycogen synthesis to be altered independently of each other and independently of changes in the rate of glycolysis. This independence allows a greater latitude of response for the individual pathways in more severe metabolic stress or in accomodating the metabolic changes necessary for long-term survival.

L8 ANSWER 45 OF 59 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.  
AN 1984:350331 BIOSIS  
DN BA78:86811  
TI IDENTIFICATION OF GTP AS A PHYSIOLOGICALLY RELEVANT **INHIBITOR** OF  
ESCHERICHIA-COLI **ADP GLUCOSE**  
**SYNTHETASE** EC-2.7.7.27.  
AU DIETZLER D N; PORTER S E; ROTH W G; LECKIE M P  
CS DIV. LAB. MED., DEP. PATHOLOGY, WASHINGTON UNIV. MED. SCH., ST. LOUIS,  
MISSOURI 63110.  
SO BIOCHEM BIOPHYS RES COMMUN, (1984) 122 (1), 289-296.  
CODEN: BBRCA9. ISSN: 0006-291X.  
FS BA; OLD  
LA English  
AB Physiological concentrations of GTP significantly **inhibited**  
wild-type E. **coli ADP-glucose**  
**synthetase** (the rate-limiting enzyme of bacterial glycogen  
synthesis). Mutant-strain enzymes known to show less **inhibition**  
by physiological AMP levels also showed less **inhibition** by  
physiological levels of GTP. This decreased **inhibition** by AMP  
and GTP can almost totally account for the higher rates of glycogen  
synthesis observed in the mutant strains. In metabolic conditions where  
glycogen synthesis increases, cellular GTP levels decrease. Thus, GTP  
**inhibition** is physiologically relevant.

-- d bib ab 43 53 56 60 70 72 79 82

L5 ANSWER 43 OF 83 CA COPYRIGHT 2003 ACS  
AN 117:187205 CA  
TI Comparison of proteins of ADP-glucose pyrophosphorylase from diverse sources  
AU Smith-White, Brian J.; Preiss, Jack  
CS Dep. Biochem., Michigan State Univ., East Lansing, MI, 48824, USA  
SO Journal of Molecular Evolution (1992), 34(5), 449-64  
CODEN: JMEVAU; ISSN: 0022-2844  
DT Journal  
LA English  
AB The primary structures of 11 proteins of ADP-glucose pyrophosphorylase are aligned and compared for relationships among them. These comparisons indicate that many domains are retained in the proteins from both the enteric bacteria and the proteins from angiosperm plants. The proteins from angiosperm plants show two main groups, with one of the main groups demonstrating two subgroups. The two main groups of angiosperm plant proteins are based upon the two subunits of the enzyme, whereas the subgroups of the large subunit group are based upon the tissue in which the particular gene had been expressed. Addnl., the small subunit group shows a slight but distinct division into a grouping based upon whether the protein is from a monocot or dicot source. Previous structure-function studies with the Escherichia coli enzyme have identified regions of the primary structure assocd. with the substrate binding site, the allosteric activator binding site, and the allosteric **inhibitor** binding site. There is conservation of the primary structure of the polypeptides for the substrate binding site and the allosteric activator binding site. The nucleotide sequences of the coding regions of the genes of 11 of these proteins are compared for relationships among them. This anal. indicates that the protein for the small subunit has been subject to greater selective pressure to retain a particular primary structure. Also, the coding region of the precursor gene for the small subunit diverged from the coding region of the precursor gene for the large subunits slightly prior to the divergence of the two coding regions of the genes for the two tissue-specific large subunit genes.

L5 ANSWER 53 OF 83 CA COPYRIGHT 2003 ACS  
AN 106:29265 CA  
TI Identification of the Escherichia coli ADP-glucose synthetase **inhibitor** binding site(s)  
AU Larsen, Charles Eric  
CS Univ. California, Davis, CA, USA  
SO (1986) 148 pp. Avail.: Univ. Microfilms Int., Order No. DA8621503  
From: Diss. Abstr. Int. B 1986, 47(6), 2416  
DT Dissertation  
LA English  
AB Unavailable

L5 ANSWER 56 OF 83 CA COPYRIGHT 2003 ACS  
AN 105:56941 CA  
TI Covalent modification of the **inhibitor** binding site(s) of Escherichia coli ADP-glucose synthetase: specific incorporation of the photoaffinity analog 8-azido-adenosine 5'-monophosphate  
AU Larsen, Charles E.; Preiss, Jack  
CS Dep. Biochem. Biophys., Univ. California, Davis, CA, 95616, USA  
SO Biochemistry (1986), 25(15), 4371-6  
CODEN: BICHAW; ISSN: 0006-2960  
DT Journal  
LA English  
AB The photoaffinity agent 8-azido-AMP (8-N3AMP) is an **inhibitor** site-specific probe of the E. coli ADP-glucose synthetase (I). In the



absence of light, 8-N3AMP exhibited the typical reversible allosteric kinetics of the physiol. **inhibitor** AMP. In the presence of light (254 nm), the analog specifically and covalently modified I, and photoincorporation was linearly related to loss of catalytic activity to .gtoreq.65% inactivation. The substrate ADP-glucose provided nearly 100% protection from 8-N3AMP photoinactivation, whereas the substrate ATP provided .apprx.50% protection and AMP, .apprx.30% protection. These 3 adenylate allosteric effectors of E. coli I also protected it from photoincorporation of 8-N3AMP. A structural overlap of the **inhibitor**- and substrate-binding sites is proposed which explains the protection data in light of the known binding and kinetic properties of this tetrameric enzyme.

- L5 ANSWER 60 OF 83 CA COPYRIGHT 2003 ACS  
 AN 101:106377 CA  
 TI Identification of GTP as a physiologically relevant **inhibitor** of Escherichia coli ADP-glucose synthetase  
 AU Dietzler, David N.; Porter, Sharon E.; Roth, William G.; Leckie, Mary P.  
 CS Sch. Med., Washington Univ., St. Louis, MO, 63110, USA  
 SO Biochemical and Biophysical Research Communications (1984), 122(1), 289-96  
 CODEN: BBRCA9; ISSN: 0006-291X  
 DT Journal  
 LA English  
 AB Physiol. concns. of GTP can significantly inhibit wild-type E. coli ADP-glucose synthetase (the rate-limiting enzyme of bacterial glycogen synthesis); mutant-strain enzymes, known to show less inhibition by physiol. AMP levels, also show less inhibition by physiol. levels of GTP. This decreased inhibition by both AMP and GTP can almost totally account for the higher cellular rates of glycogen synthesis obsd. in the mutant strains. In addn., in metabolic conditions where cellular glycogen synthesis increases, cellular GTP levels are known to decrease. Thus, GTP inhibition is physiol. relevant.
- L5 ANSWER 70 OF 83 CA COPYRIGHT 2003 ACS  
 AN 91:171367 CA  
 TI Evidence for new factors in the coordinate regulation of energy metabolism in Escherichia coli. Effects of hypoxia, chloramphenicol succinate, and 2,4-dinitrophenol on glucose use, glycogen synthesis, adenylate energy charge, and hexose phosphates in the first two periods of nitrogen starvation  
 AU Dietzler, David N.; Leckie, Mary P.; Lewis, John W.; Porter, Sharon E.; Taxman, Thomas L.; Lais, Cynthia J.  
 CS Sch. Med., Washington Univ., St. Louis, MO, 63110, USA  
 SO Journal of Biological Chemistry (1979), 254(17), 8295-307  
 CODEN: JBCHA3; ISSN: 0021-9258  
 DT Journal  
 LA English  
 AB The effects of decreased aeration, chloramphenicol succinate, and 2,4-dinitrophenol on the cellular rates of glycogen synthesis and glucose utilization and on the cellular concns. of adenine nucleotides, glucose 6-phosphate, fructose 1,6-diphosphate, and phosphoenolpyruvate during the first 2 periods of N starvation of E. coli W4597(K) were studied. A quant. relation between the changes in the rates and the accompanying changes in the hexose phosphates is demonstrated. However, the relation for glycogen synthesis is different in different sets of metabolic conditions. It is suggested that this difference reflects a change in the steady state level of a previously unknown effector of ADP-glucose synthetase (EC 2.7.7.27), the rate-limiting enzyme of bacterial glycogen synthesis. The properties of the hypothetical in vivo effector were consistent with the inhibitory effects of ppGpp (guanosine 3'-diphosphate 5'-diphosphate) and pppGpp (guanosine 3'--diphosphate 5'-triphosphate) on this enzyme in vitro. In addn., tetracycline, an **inhibitor** of the synthesis of these nucleotides, apparently prevents the change in the quant. relation. The relation between glucose utilization and the hexose

phosphates is altered at the transition to Period II of N starvation. In contrast to the ATP-hexose phosphate system of shared regulatory effects, the specific effects of the unknown effectors allow the rates of glucose utilization and glycogen synthesis to be altered independently of each other and independently of changes in the rate of glycolysis. This independence allows a greater latitude of response for the individual pathways in more severe metabolic stress or in accommodating the metabolic changes necessary for long term survival.

- L5 ANSWER 72 OF 83 CA COPYRIGHT 2003 ACS  
AN 89:175690 CA  
TI Biosynthesis of bacterial glycogen. Incorporation of pyridoxal phosphate into the allosteric activator site and an ADP-glucose-protected pyridoxal phosphate binding site of Escherichia coli B ADP-glucose synthase  
AU Parsons, Thomas F.; Preiss, Jack  
CS Dep. Biochem. Biophys., Univ. California, Davis, CA, USA  
SO Journal of Biological Chemistry (1978), 253(17), 6197-202  
CODEN: JBCHA3; ISSN: 0021-9258  
DT Journal  
LA English  
AB Pyridoxal phosphate-3H (I-3H) was covalently incorporated into E. coli B mutant strain AC70R1 ADP-glucose synthase by redn. with NaBH<sub>4</sub>. Two distinct lysine residues can be modified by the allosteric activator, I. The presence of ADP-glucose + MgCl<sub>2</sub> prevented pyridoxylation of an ADP-glucose-protected site and allowed modification of the allosteric activator site. The presence of the allosteric effectors fructose diphosphate, AMP, or hexanediol-1,6-diphosphate protected against pyridoxylation of the allosteric activator site and allowed modification of the ADP-glucose-protected site. Incorporation of I into the allosteric activator site gave modified enzyme in a high activity form, even in the absence of fructose diphosphate. This modified enzyme, when assayed in the absence of fructose diphosphate, exhibited activation kinetics similar to nonpyridoxylated enzyme assayed in the presence of fructose diphosphate and was still inhibited by AMP. These data suggest that the allosteric activator site of pyridoxylation is the fructose diphosphate binding site, and is distinct from the **inhibitor** AMP binding site. Incorporation of I into the ADP-glucose-protected site gave a decrease in enzyme activity. This pyridoxylated lysine could be involved in the binding of the substrates ADP-glucose, .alpha.-glucose 1-phosphate, or inorg. pyrophosphate, or in the catalytic mechanism of the enzyme.
- L5 ANSWER 79 OF 83 CA COPYRIGHT 2003 ACS  
AN 70:25822 CA  
TI Activator-**inhibitor**-metal ion interrelations of the adenosine diphospho-glucose pyrophosphorylase from Escherichia coli B  
AU Gentner, Norman E.  
CS Univ. of California, Davis, CA, USA  
SO (1968) 167 pp. Avail.: 68-14,596  
From: Diss. Abstr. B 1968, 29(4), 1256-7  
DT Dissertation  
LA English  
AB Unavailable
- L5 ANSWER 82 OF 83 CA COPYRIGHT 2003 ACS  
AN 67:18058 CA  
TI Activator-**inhibitor** interactions in the adenosinediphosphate glucose pyrophosphorylase of Escherichia coli B  
AU Gentner, Norman; Preiss, Jack  
CS Univ. of California, Davis, CA, USA  
SO Biochemical and Biophysical Research Communications (1967), 27(3), 417-23  
CODEN: BBRCA9; ISSN: 0006-291X  
DT Journal  
LA English  
AB The concn. of an activator of the title enzyme (I) fructose-1,6-

diphosphate (II), modulated the sensitivity of the rate of synthesis of adenosinediphosphate glucose (ADP-glucose) to inhibition by AMP and inorganic phosphate in *E. coli* B. Subsatg. levels of II-sensitized I to AMP inhibition. At low II levels, or in its absence, the rate of synthesis was relatively insensitive to inhibition by AMP. The binding of 1 mol. of **inhibitor** appeared to hinder the binding of another mol. of **inhibitor**, instead of facilitating it. The concns. of ATP and MgCl<sub>2</sub> might also be involved in the reversal of AMP inhibition. II activated I by increasing the V<sub>max</sub>. and by increasing the affinity of the enzyme for its substrates.

=> d his

(FILE 'HOME' ENTERED AT 09:53:31 ON 12 MAY 2003)

FILE 'CA' ENTERED AT 10:00:15 ON 12 MAY 2003

L1 0 S PD 1981  
L2 50 S LECKIE M?/AU  
L3 0 S S PD=1981  
L4 0 S PD=1081  
L5 508050 S PD=1981  
L6 2 S L5 AND L2  
L7 17 S PREISS J/AU  
L8 418 S PREISS J?/AU  
L9 516204 S PD=1983  
L10 9 S L8 AND L9  
L11 36 S ADP-GLUCOSE SYNTHETASE  
L12 653062 S INHIBITOR  
L13 11 S L11 AND L12

=> log hold

COST IN U.S. DOLLARS	SINCE FILE	TOTAL
	ENTRY	SESSION
FULL ESTIMATED COST	49.94	50.57
DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)	SINCE FILE	TOTAL
	ENTRY	SESSION
CA SUBSCRIBER PRICE	-4.34	-4.34

SESSION WILL BE HELD FOR 60 MINUTES  
STN INTERNATIONAL SESSION SUSPENDED AT 10:11:18 ON 12 MAY 2003

=> d his

(FILE 'HOME' ENTERED AT 16:59:22 ON 08 MAY 2003)

FILE 'CA' ENTERED AT 16:59:32 ON 08 MAY 2003

L1 FILE 'REGISTRY' ENTERED AT 16:59:37 ON 08 MAY 2003  
1 S E.C. 2.7.7.27/CN

FILE 'CA' ENTERED AT 17:00:35 ON 08 MAY 2003  
S 9027-71-8/REG#

L2 FILE 'REGISTRY' ENTERED AT 17:01:00 ON 08 MAY 2003  
1 S 9027-71-8/RN

L3 FILE 'CA' ENTERED AT 17:01:00 ON 08 MAY 2003  
733 S L2  
L4 652233 S INHIBITOR  
L5 83 S L3 AND L4

=> log hold

COST IN U.S. DOLLARS

SINCE FILE	TOTAL
ENTRY	SESSION
47.07	55.18

FULL ESTIMATED COST

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)

SINCE FILE	TOTAL
ENTRY	SESSION
-3.72	-3.72

CA SUBSCRIBER PRICE

SESSION WILL BE HELD FOR 60 MINUTES  
STN INTERNATIONAL SESSION SUSPENDED AT 17:06:53 ON 08 MAY 2003

=> d his

(FILE 'HOME' ENTERED AT 09:20:32 ON 09 MAY 2003)

FILE 'BIOSIS' ENTERED AT 09:20:53 ON 09 MAY 2003

L1 526 S ADP GLUCOSE PYROPHOSPHORYLASE  
L2 48 S ADP GLUCOSE SYNTHETASE  
L3 10 S ADP GLUCOSE SYNTHASE  
L4 584 S L1 OR L2 OR L3  
L5 1112601 S INHIBIT?  
L6 129 S L4 AND L5  
L7 760659 S BACTERIA OR COLI  
L8 59 S L7 AND L6

FILE 'WPIDS' ENTERED AT 09:33:31 ON 09 MAY 2003

L9 0 S L4 AND L5 AND L7  
L10 44 S L4

FILE 'USPATFULL' ENTERED AT 09:36:01 ON 09 MAY 2003

L11 178 S L4  
L12 407271 S L5  
L13 113703 S L7  
L14 161 S L11 AND L12 AND L13  
L15 42 S L13 (P) L11  
L16 407271 S L5  
L17 1 S 2001:121578/AN  
L18 1 S L16 AND L17

=> log hold

COST IN U.S. DOLLARS

SINCE FILE

TOTAL

ENTRY

SESSION

FULL ESTIMATED COST

14.95

58.80

SESSION WILL BE HELD FOR 60 MINUTES

STN INTERNATIONAL SESSION SUSPENDED AT 09:41:34 ON 09 MAY 2003